

[illegible]

FIGURE 3

SEQ ID NO 1: NH₂-CSESDPRGVTSYLSPPSPLDLYVHKAPKIT-COOH

FIGURE 4

SEQ ID NO 2 : NH₂-CLVVDLATMEGMNLTWYRESKEPVNPGPLNK-COOH

FIGURE 5

SEQ ID NO 3: NH₂-KDFHNGTITVTSTLPVNTNDWIEGETYY-COOH

FIGURE 6

SEQ ID NO 4: NH₂-CRVTHPHLPKDIVRSIAKAPGKRAP-COOH

FIGURE 7

SEQ ID NO 5: NH₂-LSPPSPLDLYVHKAPKITCLVVDLATME-COOH

FIGURE 8

SEQ ID NO 6:

NH₂-CGMNLTWYRESKEPVNPGPLNKKDFHNGTITVTS-COOH

FIGURE 9

SEQ ID NO 7: NH₂-TLPVNTNDWIEGETYYCRVTHPHLPK-COOH

FIGURE 10

SEQ ID NO 8: NH₂-CADSNPRGV SAYLSRPSPFDLFIRKSPTIT-COOH

FIGURE 11

SEQ ID NO 9:

NH₂-CLVVDLAPSKGTVNLTWSRASGKPVNHSTRKEE-COOH

FIGURE 12

SEQ ID NO: 10: NH₂-KQRNGTLTVTSTLPVGTRDWIEGETYQ-COOH

FIGURE 13

SEQ ID NO: 11: NH₂-CRVTHPHLPRALMRSTTKTSGPRAAP-COOH

FIGURE 14

SEQ ID 12: NH₂-SRPSPFDLFIRKSPTITCLVVDLAPSK-COOH

FIGURE 15

SEQ ID NO: 13:

NH₂-GTVNLTWSRASGKPVNHSTRKEEKQRNGTLTVTS-COOH

FIGURE 16

SEQ ID NO: 14: NH₂-TLPVGTRDWIEGETYQCRVTHPHLPR-COOH

FIGURE 17

SEQ ID NO 15:

TGCTCTGACCCGCGTGGTGTACCTCTTACCTGTCTCCGCCGTCTCCGCTGGAC
CTGTACGTTACAAAGCTCCGAAAATCACC

FIGURE 18

SEQ ID NO 16:

TGCCTGGTAGTGGACCTGGCCACCATGGAAGGCATGAACCTGACCTGGTACCG
GGAGAGCAAAGAACCCGTGAACCCGGGCCCTTTGAACAAG

FIGURE 19

SEQ ID NO 17:

TGCAAGGATCACTTCAATGGGACGATCACAGTCACGTCTACCCTGCCAGTGAAC
ACCAATGACTGGATCGAGGGCGAGACCTACTAT

FIGURE 20

SEQ ID NO 18:

TGCAGGGTGACCCACCCGCACCTGCCCAAGGACATCGTGCGCTCCATTGCCAA
GGCCCCTGGTAAGCGTGCCCCC

FIGURE 21

SEQ ID NO 19:

CTGTCTCCGCCGTCTCCGCTGGACCTGTACGTTACAAAGCTCCGAAAATCACC
TGCCTGGTAGTGGACCTGGCCACCATGGAA

FIGURE 22

SEQ ID NO 20:

TGCGGCATGAACCTGACCTGGTACCGGGAGAGCAAAGAACCCGTGAACCCGG
GCCCTTTGAACAAGAAGGATCACTTCAATGGGACGATCACAGTCACGTCT

FIGURE 23

SEQ ID NO 21:

ACCCTGCCAGTGAACACCAATGACTGGATCGAGGGCGAGACCTACTATTGCAG
GGTGACCCACCCGCACCTGCCCAAG

FIGURE 24

SEQ ID NO 22:

TGCGCGGACAGCAACCCGAGAGGGGTGAGCGCCTACCTAAGCCGGCCCAGCC
CGTTCGACCTGTTTCATCCGCAAGTCGCCCCACGATCACC

FIGURE 25

SEQ ID NO 23:

TGTCTGGTGGTGGACCTGGCACCCAGCAAGGGGACCGTGAACCTGACCTGGTC
CCGGGCCAGTGGGAAGCCTGTGAACCACTCCACCAGAAAGGAGGAG

FIGURE 26

SEQ ID NO 24:

AAGCAGCGCAATGGCACGTTAACCGTCACGTCCACCCTGCCGGTGGGCACCCG
AGACTGGATCGAGGGGGAGACCTACCAG

FIGURE 27

SEQ ID NO 25:

TGCAGGGTGACCCACCCCCACCTGCCCAGGGCCCTCATGCGGTCCACGACCA
AGACCAGCGGCCCCGCGTGCTGCCCCG

FIGURE 28

SEQ ID NO 26:

AGCCGGCCCAGCCCGTTCGACCTGTTTCATCCGCAAGTCGCCCCACGATCACCTG
TCTGGTGGTGGACCTGGCACCCAGCAAG

FIGURE 29

SEQ ID NO 27:

GGGACCGTGAACCTGACCTGGTCCCGGGCCAGTGGGAAGCCTGTGAACCACT
CCACCAGAAAGGAGGAGAAGCAGCGCAATGGCACGTTAACCGTCACGTCC

FIGURE 30

SEQ ID NO 28:

ACCCTGCCGGTGGGCACCCGAGACTGGATCGAGGGGGAGACCTACCAGTGCA
GGGTGACCCACCCCCACCTGCCCAGG